

1/5

Is 26	1	MRHMQSVTRASSI	CGSGVKQVI	GKGHPHARV	VGARAKA	QIPEREE
Is 27	1	MRHMQSATRATL	VCGSGVKQII	AKGHPNTRV	FGARAKA	QIPEREV
Is 26	45	LSVKPKMVRNTH	LNLPQERQAF	YRLL	ENEQIQEFL	SMDSCLRI
Is 27	45	LAAKPKITRITH	LNLPQERQAF	YRLL	ENEQIQEFL	SMDSCLKI
Is 26	89	SDKYLIAMVLA	YFKRAAGLYT	SEYTTMNF	FFVALYL	ANDMEEDEE
Is 27	89	SDKYLIAMVLA	YFKRA - GLYT	SEYTTMNF	FFVALYL	ANDMEEDEE
Is 26	133	DYKYEI	FPWALGDSWREL	FPQFLRL	RDDFWAKMNY	RAVVSRRCC
Is 27	132	DYKYEI	FPWALGDSWREF	FPQFLRL	RDDFWAKMNY	RAVVSRRCC
Is 26	177	DEVMSKDPTH	HWAWLRDRP	MHHSGAMRGYL	RNEDDFF	PRGPGGLTP
Is 27	176	DEVMAKDPTH	HWAWLRDRPI	HHSGALRGYL	RNEDDFF	PRGPGGLTP
Is	221	ASCTLCHKAGV	CDSGGVSHNNSS	SPEQEI	FHYTNREWSQ	ELLML
Is 27	220	ASCALCHKASV	CDSGGVSHDNSS -	PEQEI	FHYTNREWSQ	ELLIL
Is 26	265	PPELLLDPECT	HDLHI	LQEPLV	GLEPDGTAL	EWHL
Is 27	263	PPELLLDPESTY	DIHI	FQEPLV	GLEPDGAAL	EWHL

Fig.1

2/5

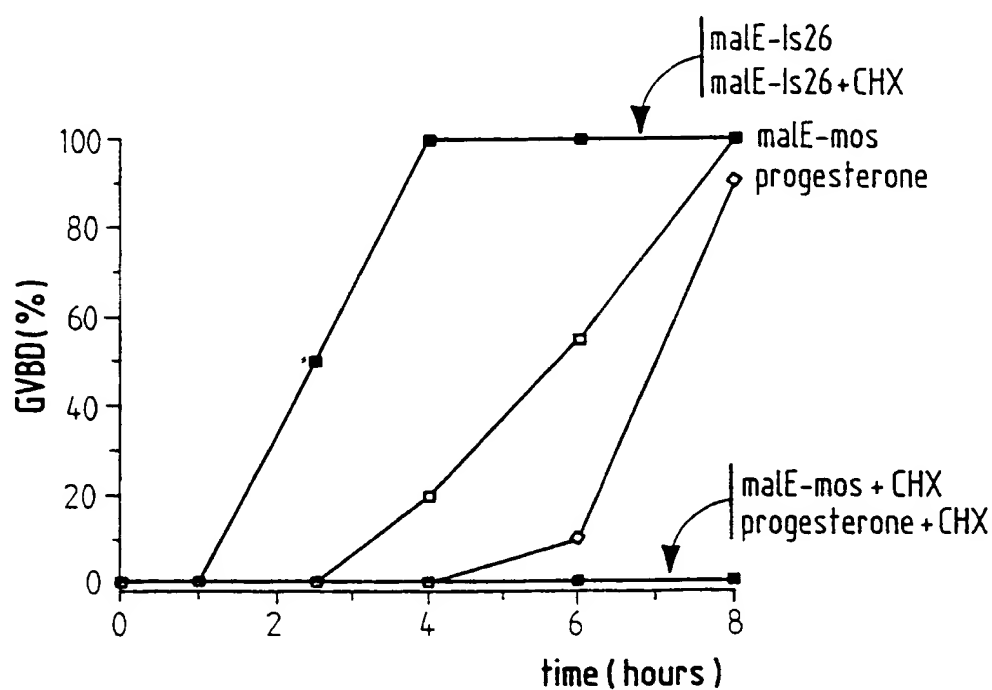


Fig. 2

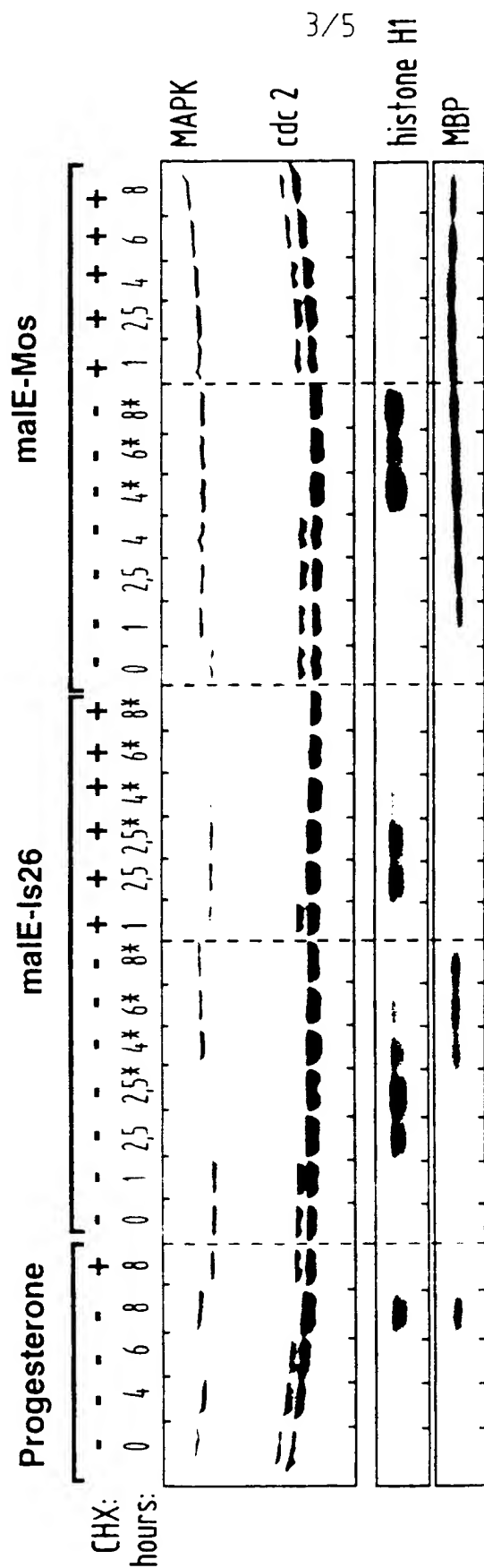


Fig. 3

4/5

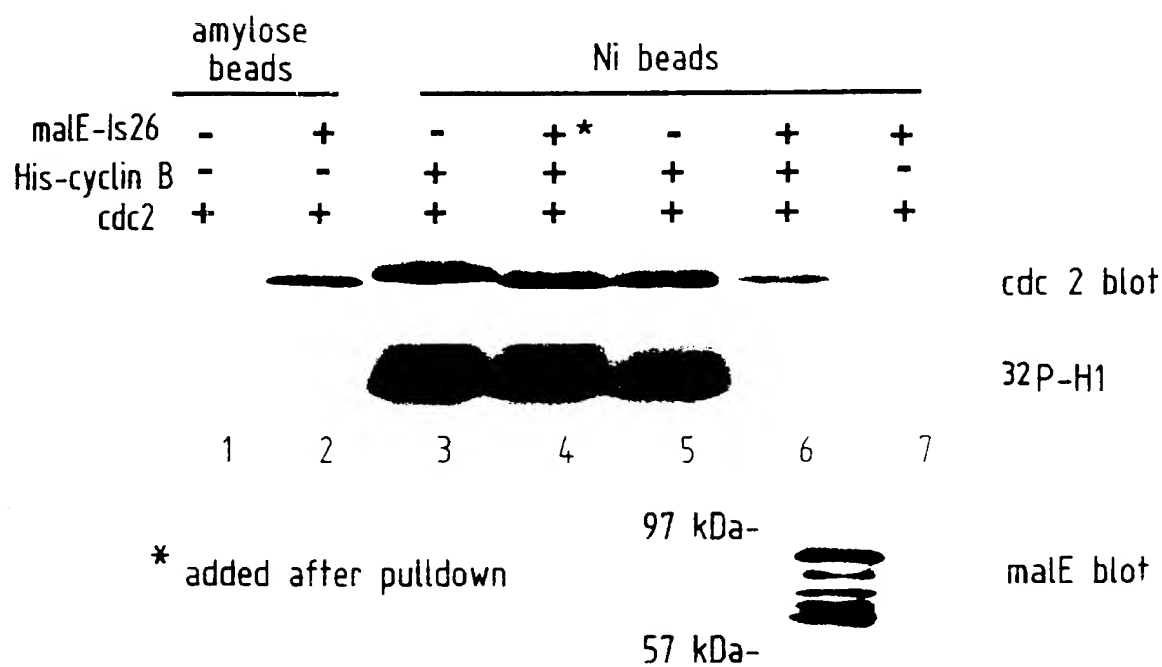


Fig. 4

5/5

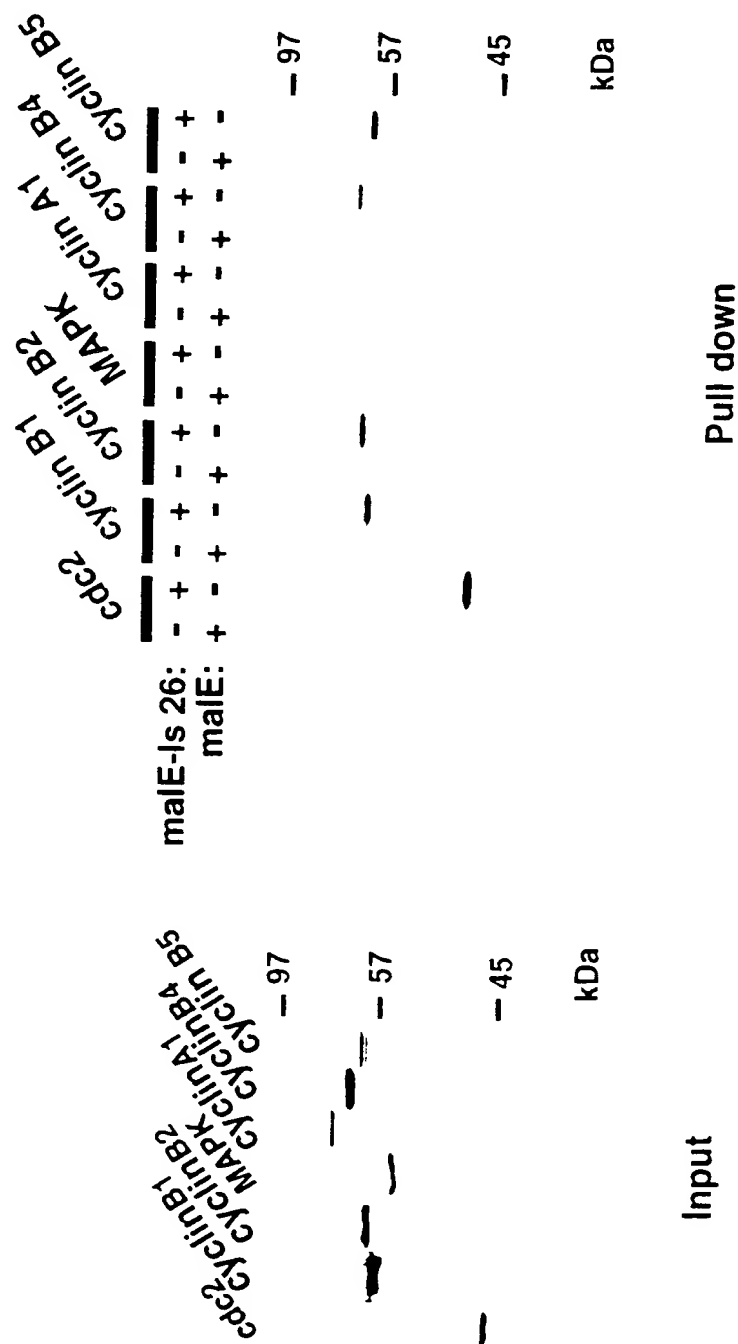


Fig. 5